

RAW SEQUENCE LISTING

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Application Serial Number: 091674,237C
Source: FW16
Date Processed by STIC: 5-25-05

ENTERED



IFW16

RAW SEQUENCE LISTING

DATE: 05/25/2005

PATENT APPLICATION: US/09/674,237C

TIME: 13:53:57

Input Set : A:\3477.89.ST25.txt

Output Set: N:\CRF4\05252005\I674237C.raw

(ps.c)

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3 <110> APPLICANT: Egan, Sean E.
4      Wang, Wei
5      Sengar, Ameet
7 <120> TITLE OF INVENTION: ESE GENES AND PROTEINS
9 <130> FILE REFERENCE: 3477-89
11 <140> CURRENT APPLICATION NUMBER: US 09/674,237C
12 <141> CURRENT FILING DATE: 2001-07-24
14 <150> PRIOR APPLICATION NUMBER: PCT/CA99/00375
15 <151> PRIOR FILING DATE: 1999-04-27
17 <150> PRIOR APPLICATION NUMBER: US 60/118,739
18 <151> PRIOR FILING DATE: 1999-02-05
20 <150> PRIOR APPLICATION NUMBER: CA 2230201
21 <151> PRIOR FILING DATE: 1998-04-27
23 <160> NUMBER OF SEQ ID NOS: 37
25 <170> SOFTWARE: PatentIn version 3.3
27 <210> SEQ ID NO: 1
28 <211> LENGTH: 5084
29 <212> TYPE: DNA
30 <213> ORGANISM: Mus musculus
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213 <223> OTHER INFORMATION: Mouse Esei
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218 1 5 10 15
220 ata act gtg gag gaa agg gcc aag cat gac cag cag ttc ctt agc ctg 96
221 Ile Thr Val Glu Glu Arg Ala Lys His Asp Gln Gln Phe Leu Ser Leu
222 20 25 30
224 aag ccg ata gcg gga ttt att act ggt gat caa gcg agg aac ttt ttt 144
225 Lys Pro Ile Ala Gly Phe Ile Thr Gly Asp Gln Ala Arg Asn Phe Phe
226 35 40 45
228 ttc caa tct ggg tta cct cag cct gtc tta gca caa ata tgg gcg cta 192
229 Phe Gln Ser Gly Leu Pro Gln Pro Val Leu Ala Gln Ile Trp Ala Leu
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232 gcg gac atg aat aac gat gga agg atg gat caa gtg gaa ttt tcc ata 240
233 Ala Asp Met Asn Asn Asp Gly Arg Met Asp Gln Val Glu Phe Ser Ile
234 65 70 75 80
236 gcc atg aag ctt atc aaa ctg aag cta caa gga tat cag ctc ccc tcc 288
237 Ala Met Lys Leu Ile Lys Leu Lys Leu Gly Tyr Gln Leu Pro Ser
238 85 90 95
240 aca ctt ccc cct gtc atg aaa cag caa cca gtg gct att tcc agt gca 336
241 Thr Leu Pro Pro Val Met Lys Gln Gln Pro Val Ala Ile Ser Ser Ala
242 100 105 110
244 cca gca ttt ggt ata gga ggg att gct agc atg cca cca ctc aca gct 384
245 Pro Ala Phe Gly Ile Gly Gly Ile Ala Ser Met Pro Pro Leu Thr Ala
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248 gtt gct cct gtg cca atg ggc tcc att cca gtt gtt gga atg tct cca 432
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256	ggg	gct	cct	ccc	gtc	ata	cag	cct	ctg	cct	gcg	ttt	gcg	cat	cct	gca	528
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264	caa	tta	aac	act	aag	tta	cag	aag	gca	caa	tca	ttc	gat	gtc	gcc	agc	624
265	Gln	Leu	Asn	Thr	Lys	Leu	Gln	Lys	Ala	Gln	Ser	Phe	Asp	Val	Ala	Ser	
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268	gcc	cct	cca	gca	gca	gaa	tgg	gct	gtg	cct	cag	tca	tca	agg	ctg	aaa	672
269	Ala	Pro	Pro	Ala	Ala	Glu	Trp	Ala	Val	Pro	Gln	Ser	Ser	Arg	Leu	Lys	
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276	aca	ggt	ccc	cag	gca	aga	act	att	ctc	atg	caa	tca	agt	tta	ccc	cag	768
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282			260						265					270			
284	aaa	ctc	act	gca	gaa	gaa	ttt	atc	cta	gct	atg	cac	cta	att	gat	gtt	864
285	Lys	Leu	Thr	Ala	Glu	Glu	Phe	Ile	Leu	Ala	Met	His	Leu	Ile	Asp	Val	
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292	cct	cct	tcc	ttc	aga	aga	gtt	cgc	tcc	ggc	agt	ggg	atg	tcc	gtc	ata	960
293	Pro	Pro	Ser	Phe	Arg	Arg	Val	Arg	Ser	Gly	Ser	Gly	Met	Ser	Val	Ile	
294	305				310					315					320		
296	agc	tct	tct	tct	gtg	gat	cag	agg	ctg	cct	gag	gag	ccg	tcg	tca	gag	1008
297	Ser	Ser	Ser	Ser	Val	Asp	Gln	Arg	Leu	Pro	Glu	Glu	Pro	Ser	Ser	Glu	
298				325						330					335		
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301	Asp	Glu	Gln	Gln	Pro	Glu	Lys	Lys	Leu	Pro	Val	Thr	Phe	Glu	Asp	Lys	
302			340						345					350			
304	aag	cgg	gag	aac	ttc	gag	cga	ggc	agt	gtg	gag	ctg	gag	aag	cgc	cgc	1104
305	Lys	Arg	Glu	Asn	Phe	Glu	Arg	Gly	Ser	Val	Glu	Leu	Glu	Lys	Arg	Arg	
306			355					360					365				
308	caa	gcg	ctc	ttg	gag	cag	cag	cgc	aaa	gag	cag	gag	cgg	ttg	gct	cag	1152
309	Gln	Ala	Leu	Leu	Glu	Gln	Gln	Arg	Lys	Glu	Gln	Glu	Arg	Leu	Ala	Gln	
310		370				375						380					
312	ctg	gag	cgc	gcc	gag	cag	gag	agg	aaa	gag	cgg	gag	cgc	cag	gag	cag	1200
313	Leu	Glu	Arg	Ala	Glu	Gln	Glu	Arg	Lys	Glu	Arg	Glu	Arg	Gln	Glu	Gln	
314	385				390					395					400		
316	gag	gcc	aag	cgg	cag	ctg	gag	ctg	gag	aag	cag	ctg	gag	aag	cag	cgg	1248

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321	Glu	Leu	Glu	Arg	Gln	Arg	Glu	Glu	Arg	Arg	Lys	Glu	Ile	Glu	Arg		
322				420					425				430				
324	cgc	gag	gcc	gca	aaa	cgg	gaa	ctg	gaa	agg	cag	cga	caa	ctt	gaa	tgg	1344
325	Arg	Glu	Ala	Ala	Lys	Arg	Glu	Leu	Glu	Arg	Gln	Arg	Gln	Leu	Glu	Trp	
326			435				440					445					
328	gaa	cgg	aac	cgg	aga	cag	gaa	ctc	ctg	aat	cag	agg	aac	aag	gag	cag	1392
329	Glu	Arg	Asn	Arg	Arg	Gln	Glu	Leu	Leu	Asn	Gln	Arg	Asn	Lys	Glu	Gln	
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332	gag	ggc	acc	gtg	gtc	ctg	aag	gca	agg	agg	aag	act	ctg	gag	ttt	gag	1440
333	Glu	Gly	Thr	Val	Val	Leu	Lys	Ala	Arg	Arg	Lys	Thr	Leu	Glu	Phe	Glu	
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337	Leu	Glu	Ala	Leu	Asn	Asp	Lys	Lys	His	Gln	Leu	Glu	Gly	Lys	Leu	Gln	
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353	Gln	Ile	Leu	Ser	Asp	Gln	Leu	Lys	Gln	Val	Gln	Gln	Asn	Ser	Leu	His	
354	545				550				555				560				
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369	Leu	Arg	Glu	Ile	His	Ser	Lys	Gln	Gln	Leu	Gln	Lys	Gln	Arg	Ser	Leu	
370		610				615				620							
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373	Glu	Ala	Ala	Arg	Leu	Lys	Gln	Lys	Glu	Gln	Glu	Arg	Lys	Ser	Leu	Glu	
374	625				630				635				640				
376	tta	gag	aag	caa	aag	gaa	gac	gct	cag	aga	cga	ggt	cag	gaa	agg	gac	1968
377	Leu	Glu	Lys	Gln	Lys	Glu	Asp	Ala	Gln	Arg	Arg	Val	Gln	Glu	Arg	Asp	
378			645				650			655							
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TIME: 13:53:58

Input Set : A:\3477.89.ST25.txt
Output Set: N:\CRF4\05252005\I674237C.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:18; N Pos. 102,103

Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete, per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:28,29,30,31,32,33,36,37

VERIFICATION SUMMARY

DATE: 05/25/2005

PATENT APPLICATION: US/09/674,237C

TIME: 13:53:58

Input Set : A:\3477.89.ST25.txt

Output Set: N:\CRF4\05252005\I674237C.raw

L:2218 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18 after pos.:60